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ENTERED

1 <110> APPLICANT: Chiaur, D.

OIPE

RAW SEQUENCE LISTINGPATENT APPLICATION: **US/10/652,928**DATE: 09/17/2003
TIME: 09:43:49

Input Set : N:\Crf3\RULE60\10652928.raw.txt
Output Set: N:\CRF4\09172003\J652928.raw

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Pagano, M.
      2
      3
              Latres, E.
      4 <120> TITLE OF INVENTION: NOVEL UBIQUITIN LIGASES AS THERAPEUTIC TARGETS
      5 <130> FILE REFERENCE: 5914-081
      6 <140> CURRENT APPLICATION NUMBER: 10/652,928
C--> 7 <141> CURRENT FILING DATE: 2003-08-28
      8 <150> PRIOR APPLICATION NUMBER: US/09/385,219A
      9 <151> PRIOR FILING DATE: 1999-08-27
     10 <150> PRIOR APPLICATION NUMBER: 60/098,355
     11 <151> PRIOR FILING DATE: 1998-08-28
     12 <150> PRIOR APPLICATION NUMBER: 60/118,568
     13 <151> PRIOR FILING DATE: 1999-02-03
     14 <150> PRIOR APPLICATION NUMBER: 60/124,449
     15 <151> PRIOR FILING DATE: 1999-03-15
     16 <160> NUMBER OF SEQ ID NOS: 90
     17 <170> SOFTWARE: PatentIn Ver. 2.0
     19 <210> SEO ID NO: 1
     20 <211> LENGTH: 2151
     21 <212> TYPE: DNA
     22 <213> ORGANISM: Homo sapiens
     23 <400> SEQUENCE: 1
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              tcggcgatta tggacccggc cgaggcggtg ctgcaagaga aggcactcaa gtttatgaat 120
     25
     26
              tecteagaga gagaagaetg taataatgge gaaceeeta ggaagataat accagagaag 180
     27
              aattcactta gacagacata caacagctgt gccagactct gcttaaacca agaaacagta 240
              tgtttagcaa gcactgctat gaagactgag aattgtgtgg ccaaaacaaa acttgccaat 300
     28
     29
              ggcacttcca gtatgattgt gcccaagcaa cggaaactct cagcaagcta tgaaaaggaa 360
     30
              aaggaactgt gtgtcaaata ctttgagcag tggtcagagt cagatcaagt ggaatttgtg 420
              gaacatetta tateecaaat gtgteattae caacatggge acataaaete gtatettaaa 480
     31
             cctatgttgc agagagattt cataactgct ctgccagctc ggggattgga tcatatcgct 540
     32
     33
             qaqaacattc tgtcatacct ggatgccaaa tcactatgtg ctgctgaact tgtgtgcaag 600
     34
             qaatgqtacc gagtgacctc tgatggcatg ctgtggaaga agcttatcga gagaatggtc 660
             aggacagatt ctctgtggag aggcctggca gaacgaagag gatggggaca gtatttattc 720
     35
             aaaaacaaac ctcctgacgg gaatgctcct cccaactctt tttatagagc actttatcct 780
     36
    37
             aaaattatac aagacattga gacaatagaa tctaattgga gatgtggaag acatagttta 840
             caqaqaattc actgccqaag tqaaacaagc aaaggagttt actgtttaca gtatgatgat 900
     38
    39
             cagaaaatag taagcggcct tcgagacaac acaatcaaga tctgggataa aaacacattg 960
     40
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             agagtgatca taacaggatc atcggattcc acggtcagag tgtgggatgt aaatacaggt 1080
    41
    42
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Input Set: N:\Crf3\RULE60\10652928.raw.txt
Output Set: N:\CRF4\09172003\J652928.raw

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46
47
         tacagggaca ggctggtagt gagtggctca tctgacaaca ctatcagatt atgggacata 1440
48
         gaatgtggtg catgtttacg agtgttagaa ggccatgagg aattggtgcg ttgtattcga 1500
49
         tttgataaca agaggatagt cagtggggcc tatgatggaa aaattaaagt gtgggatctt 1560
50
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         cattccggaa gagtttttcg actacagttt gatgaattcc agattgtcag tagttcacat 1680
51
52
         gatgacacaa teeteatetg ggaetteeta aatgateeag etgeecaage tgaaceecee 1740
         cgttcccctt ctcgaacata cacctacatc tccagataaa taaccataca ctgacctcat 1800
53
54
         acttgcccag gacccattaa agttgcggta tttaacgtat ctgccaatac caggatgagc 1860
55
         aacaacagta acaatcaaac tactgcccag tttccctgga ctagccgagg agcagggctt 1920
56
         tqaqactcct qttqqqacac aqttqqtctq cagtcqqccc aggacqgtct actcagcaca 1980
         actgactgct tcagtgctgc tatcagaaga tgtcttctat caattgtgaa tgattggaac 2040
57
58
         ttttaaacct cccctctct cctcctttca cctctgcacc tagttttttc ccattggttc 2100
59
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                                                                             2151
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62 <211> LENGTH: 569
63 <212> TYPE: PRT
64 <213> ORGANISM: Homo sapiens
65 <400> SEQUENCE: 2
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66
67
                                               10
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68
69
70
         Ile Ile Pro Glu Lys Asn Ser Leu Arg Gln Thr Tyr Asn Ser Cys Ala
71
                                       40
72
         Arg Leu Cys Leu Asn Gln Glu Thr Val Cys Leu Ala Ser Thr Ala Met
73
74
         Lys Thr Glu Asn Cys Val Ala Lys Thr Lys Leu Ala Asn Gly Thr Ser
75
                                                   75
76
         Ser Met Ile Val Pro Lys Gln Arg Lys Leu Ser Ala Ser Tyr Glu Lys
77
         Glu Lys Glu Leu Cys Val Lys Tyr Phe Glu Gln Trp Ser Glu Ser Asp
78
79
                     100
                                          105
                                                              110
80
         Gln Val Glu Phe Val Glu His Leu Ile Ser Gln Met Cys His Tyr Gln
81
                                                          125
82
         His Gly His Ile Asn Ser Tyr Leu Lys Pro Met Leu Gln Arg Asp Phe
83
                                  135
                                                      140
84
         Ile Thr Ala Leu Pro Ala Arg Gly Leu Asp His Ile Ala Glu Asn Ile
85
         145
                             150
                                                  155
86
         Leu Ser Tyr Leu Asp Ala Lys Ser Leu Cys Ala Ala Glu Leu Val Cys
87
                                              170
88
         Lys Glu Trp Tyr Arg Val Thr Ser Asp Gly Met Leu Trp Lys Lys Leu
89
                     180
                                          185
90
         Ile Glu Arg Met Val Arg Thr Asp Ser Leu Trp Arg Gly Leu Ala Glu
91
                                      200
92
         Arg Arg Gly Trp Gly Gln Tyr Leu Phe Lys Asn Lys Pro Pro Asp Gly
93
                                  215
                                                      220
         Asn Ala Pro Pro Asn Ser Phe Tyr Arg Ala Leu Tyr Pro Lys Ile Ile
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Input Set : N:\Crf3\RULE60\10652928.raw.txt
Output Set: N:\CRF4\09172003\J652928.raw

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225
                                                                       240
95
96
         Gln Asp Ile Glu Thr Ile Glu Ser Asn Trp Arg Cys Gly Arg His Ser
97
                         245
                                              250
98
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99
                     260
                                          265
                                                               270
100
          Leu Gln Tyr Asp Asp Gln Lys Ile Val Ser Gly Leu Arg Asp Asn Thr
101
                                       280
102
          Ile Lys Ile Trp Asp Lys Asn Thr Leu Glu Cys Lys Arg Ile Leu Thr
                                   295
103
          Gly His Thr Gly Ser Val Leu Cys Leu Gln Tyr Asp Glu Arg Val Ile
104
105
106
          Ile Thr Gly Ser Ser Asp Ser Thr Val Arg Val Trp Asp Val Asn Thr
107
                          325
                                               330
          Gly Glu Met Leu Asn Thr Leu Ile His His Cys Glu Ala Val Leu His
108
109
                                           345
          Leu Arg Phe Asn Asn Gly Met Met Val Thr Cys Ser Lys Asp Arg Ser
110
111
                                       360
          Ile Ala Val Trp Asp Met Ala Ser Pro Thr Asp Ile Thr Leu Arg Arg
112
113
                                   375
                                                        380
          Val Leu Val Gly His Arg Ala Ala Val Asn Val Val Asp Phe Asp Asp
114
115
                               390
                                                   395
          Lys Tyr Ile Val Ser Ala Ser Gly Asp Arg Thr Ile Lys Val Trp Asn
116
117
                          405
                                               410
          Thr Ser Thr Cys Glu Phe Val Arg Thr Leu Asn Gly His Lys Arg Gly
118
119
                      420
                                           425
120
          Ile Ala Cys Leu Gln Tyr Arg Asp Arg Leu Val Val Ser Gly Ser Ser
121
                                       440
          Asp Asn Thr Ile Arg Leu Trp Asp Ile Glu Cys Gly Ala Cys Leu Arg
122
123
                                   455
          Val Leu Glu Gly His Glu Glu Leu Val Arg Cys Ile Arg Phe Asp Asn
124
125
                               470
                                                   475
126
          Lys Arg Ile Val Ser Gly Ala Tyr Asp Gly Lys Ile Lys Val Trp Asp
127
                          485
                                              490
128
          Leu Val Ala Ala Leu Asp Pro Arg Ala Pro Ala Gly Thr Leu Cys Leu
129
                                           505
130
          Arg Thr Leu Val Glu His Ser Gly Arg Val Phe Arg Leu Gln Phe Asp
131
                                       520
132
          Glu Phe Gln Ile Val Ser Ser Ser His Asp Asp Thr Ile Leu Ile Trp
133
134
          Asp Phe Leu Asn Asp Pro Ala Ala Gln Ala Glu Pro Pro Arg Ser Pro
135
                              550
                                                   555
136
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137
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140 <211> LENGTH: 1476
141 <212> TYPE: DNA
142 <213> ORGANISM: Homo sapiens
143 <400> SEQUENCE: 3
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Input Set: N:\Crf3\RULE60\10652928.raw.txt
Output Set: N:\CRF4\09172003\J652928.raw

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145
146
          ctcaggcatc tctccaataa cctagagact ctcctcaagc gggacttcct caaactcctt 180
147
          cccctqqaqc tcaqttttta tttqttaaaa tqqctcqatc ctcaqacttt actcacatqc 240
148
          tgcctcgtct ctaaacagtg gaataaggtg ataagtgcct gtacagaggt gtggcagact 300
149
          gcatgtaaaa atttgggctg gcagatagat gattctgttc aggacgcttt gcactggaag 360
150
          aaggtttatt tgaaggctat tttgagaatg aagcaactgg aggaccatga agcctttgaa 420
151
          acctegteat taattggaca cagtgecaga gtgtatgeac tttactacaa agatggactt 480
152
          ctctgtacag ggtcagatga cttgtctgca aagctgtggg atgtgagcac agggcagtgc 540
153
          gtttatggca tccagaccca cacttgtgca gcggtgaagt ttgatgaaca gaagcttgtg 600
154
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          cactttcggg ggcacacggg ggcggtattt agcgtggact acaatgatga actggatatc 720
155
156
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157
          ctgaacacac tcaccqqqca cacqqaatqq qtcaccaaqq.taqttttqca gaaqtqcaaa 840
158
          qtcaaqtctc tcttqcacaq tcctqqaqac tacatcctct taaqtqcaqa caaatatqaq 900
159
          attaagattt ggccaattgg gagagaaatc aactgtaagt gcttaaagac attgtctgtc 960
          totgaggata gaagtatotg cotgoagoca agacttoatt ttgatggoaa atacattgto 1020
160
161
          tgtagttcag cacttggtct ctaccagtgg gactttgcca gttatgatat tctcagggtc 1080
          atcaagactc ctgagatagc aaacttggcc ttgcttggct ttggagatat ctttgccctg 1140
162
          ctgtttgaca accgctacct gtacatcatg gacttgcgga cagagagcct gattagtcgc 1200
163
          tgqcctctqc cagagtacag ggaatcaaag agaggctcaa gcttcctggc aggcgaacat 1260
164
165
          cctggctgaa tggactggat gggcacaatg acacgggctt ggtctttgcc accagcatgc 1320
166
          ctgaccacag tattcacctg gtgttgtgga aggagcacgg ctgacaccat gagccaccac 1380
          cgctgactga ctttgggtgc cggggctgcg ggttttgggt gcacctctgc ggcacgcgac 1440
167
168
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170 <210> SEQ ID NO: 4
171 <211> LENGTH: 422
172 <212> TYPE: PRT
173 <213> ORGANISM: Homo sapiens
174 <400> SEQUENCE: 4
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177
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178
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                                            25
179
          Ile Ser Leu Ser Gly Ala Val Gln Leu Arg His Leu Ser Asn Asn Leu
180
                                        40
          Glu Thr Leu Leu Lys Arg Asp Phe Leu Lys Leu Leu Pro Leu Glu Leu
181
182
183 ·
          Ser Phe Tyr Leu Leu Lys Trp Leu Asp Pro Gln Thr Leu Leu Thr Cys
184
          Cys Leu Val Ser Lys Gln Trp Asn Lys Val Ile Ser Ala Cys Thr Glu
185
186
                                                90
187
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188.
                                          105
189
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190
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1.91
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193
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194
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Input Set : N:\Crf3\RULE60\10652928.raw.txt
Output Set: N:\CRF4\09172003\J652928.raw

195 196		Leu	Cys	Thr	Gly	Ser 165	Asp	Asp	Leu	Ser	Ala 170	Lys	Leu	Trp	Asp	Val 175	Ser	
197 198		Thr	Gly	Gln	Cys 180	Val	Tyr	Gly	Ile	Gln 185	Thr	His	Thr	Cys	Ala 190	Ala	Val	
199 200		Lys	Phe	Asp 195		Gln	Lys	Leu	Val 200		Gly	Ser	Phe	Asp 205		Thr	Val	
201 202	•	Ala	Cys 210		Glu	Trp	Ser	Ser 215		Ala	Arg	Thr	Gln 220		Phe	Arg	Gly	
203				Gly	Ala	Val	Phe 230		Val	Asp	Tyr	Asn 235		Glu	Leu	Asp	Ile 240	
204		225 Leu	Val	Ser	Gly			Asp	Phe	Thr			Val	Trp	Ala	Leu		
206 207		Ala	Gly	Thr	_	245 Leu	Asn	Thr	Leu		250 Gly	His	Thr	Glu		255 Val	Thr	
208		Lys	Val		260 Leu	Gln	Lys	Cys	_	265 Val	Lys	Ser	Leu		270 His	Ser	Pro	
210 211		Gly	_	275 Tyr	Ile	Leu	Leu		280 Ala	Asp	Lys	Tyr		285 Ile	Lys	Ile	Trp	
212 213			290 Ile	Gly	Arg	Glu		295 Asn	Cys	Lys	Cys		300 Lys	Thr	Leu	Ser	Val 320	
214 215		305 Ser	Glu	Asp	Arg		310 Ile	Cys	Leu	Gln	Pro 330	315 Arg	Leu	His	Phe	Asp 335		
216 217		Lys	Tyr	Ile	Val 340	325 Cys	Ser	Ser	Ala	Leu 345		Leu	Tyr	Gln	Trp 350	Asp	Phe	
218 219		Ala	Ser	_		Ile	Leu	Arg			Lys	Thr	Pro			Ala	Asn	
220 221		Leu		355 Leu	Leu	Gly	Phe	_	360 Asp	Ile	Phe	Ala		365 Leu	Phe	Asp	Asn	
222 223			370 Tyr	Leu	Tyr	Ile		375 Asp	Leu	Arg	Thr		380 Ser	Leu	Ile	Ser		
224 225		385 Trp	Pro	Leu	Pro	Glu	390 Tyr	Arq	Glu	Ser	Lys	395 Arg	Gly	Ser	Ser	Phe	400 Leu	
226		_				405	_	_			410	,	-			415		
227 228			_		420	Pro	Gly									•		
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240																	ttcaa	
241																	cagctt	
242																	caactg	
243																	tgtgt	
244		tcgt	aaac	ctc o	caaat	ccct	g to	ette	jctta	a aga	ataga	atga	tact	ccaç	gta 🤄	gatga	tccat	600
																	•	

Input Set: N:\Crf3\RULE60\10652928.raw.txt
Output Set: N:\CRF4\09172003\J652928.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:26; Xaa Pos. 218,556,630
Seq#:29; N Pos. 13,47,68,88,270
Seq#:30; Xaa Pos. 15,22,28,89
Seq#:37; N Pos. 45,329,332
Seq#:38; Xaa Pos. 110,111
Seq#:51; N Pos. 1733
Seq#:52; Xaa Pos. 576,586
Seq#:53; N Pos. 348
Seq#:54; Xaa Pos. 150,309,340,374
Seq#:59; N Pos. 471

VERIFICATION SUMMARY

DATE: 09/17/2003

PATENT APPLICATION: US/10/652,928

TIME: 09:43:50

Input Set : N:\Crf3\RULE60\10652928.raw.txt Output Set: N:\CRF4\09172003\J652928.raw

L:7 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:964 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:208 M:341 Repeated in SeqNo=26 L:1188 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0 M:341 Repeated in SeqNo=29 L:1203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0 M:341 Repeated in SeqNo=30 L:1374 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0 M:341 Repeated in SeqNo=37 L:1403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:96 L:1948 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:1680 L:2030 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:560 M:341 Repeated in SeqNo=52 L:2049 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53 after pos.:300 L:2101 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54 after pos.:144 M:341 Repeated in SeqNo=54 L:2349 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:420

STATISTICS SUMMARY

DATE: 09/17/2003

PATENT APPLICATION: US/10/652,928

TIME: 09:43:50

Input Set : N:\Crf3\RULE60\10652928.raw.txt Output Set: N:\CRF4\09172003\J652928.raw

Application Serial Number: US/10/652,928

Alpha or Numeric or Xml: Numeric

Application Class:

Application File Date: 08-28-2003

Art Unit: OIPE

Software Application: PatentIN2.0 Total Number of Sequences: 90

Total Nucleotides: 38541 Total Amino Acids: 11248

Number of Errors: 0 Number of Warnings: 21 Number of Corrections: 1

MESSAGE SUMMARY

271 C: 1 (Current Filing Date differs) **341 W: 21** ((46) "n" or "Xaa" used)